## Amendments to the Claims:

The following listing of claims replaces all previous listings or versions thereof:

- 1-13. (Canceled).
- 14. (Currently amended) A purified human alpha subunit of an SCN1A sodium channel nucleic acid sequence comprising a nucleic acid sequence selected from the group consisting of:
  - (a) the nucleic acid sequence of SEQ ID NO:1;
  - (b) a full complement of (a);
  - [[(c)]](a) a nucleic acid sequence encoding an alpha subunit of SCN1A selected from the group consisting of:
    - (i) the alpha subunit of SCN1A set forth in SEQ ID NO:3;
    - (ii)—an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 188 which replaces an aspartic acid residue by a valine residue;
    - [[(iii)]](ii) an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 1238 which replaces a glutamic acid residue by an aspartic acid residue;
    - [[(iv)]](iii) an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 1773 which replaces a serine residue residue by a tyrosine residue; and [[v]](iv) an alpha subunit of SCN1A being at least 95% identical to the SCN1A alpha subunits in [[(ii)-(iv)]](i)-(iii) and comprising one of the
    - mutations at amino acid position 188, 1238 or 1773; [[and]]

      (b) an SCN1A nucleic acid fragment selected from the group consisting of:

      [[(vi)]](v) \_\_\_\_ an amplified segment comprising of the nucleic
      - [[(vii)]](vi) an amplified segment comprising the nucleic acid sequence from nucleotide 739 to 867 of SEQ ID NO:1 having a mutation at nucleotide 828.

acid sequence from nucleotide 739 to 867 of SEQ ID NO:1,

[[(d)]](b)

- [[(viii)]](vii) an amplified segment eomprising of the nucleic acid sequence from nucleotide 3970 to 4143 of SEO ID NO:1.
- [[(ix)]](viii) an amplified segment comprising the nucleic acid sequence from nucleotide 3970 to 4143 of SEQ ID NO:1 having a mutation at position 3978,
- [[(x)]](ix) an amplified segment emprising consisting of the nucleic acid sequence from nucleotide 5521 to 5747 of SEQ ID NO:1, and
- [[(xi)]](x) an amplified segment comprising the nucleic acid sequence from nucleotide 5521 to 5747 of SEQ ID NO:1 having a mutation at position 5582; and
- (c) a full-length complement of (a) or (b).

## 15.-16. (Canceled)

17. (Currently amended) A vector comprising any one of the sequences <u>nucleic acids</u> of claim 14.

## 18.-19. (Canceled)

20. (Previously presented) An isolated cell harboring a vector of claim 17.

## 21.-24. (Canceled)

- 25. (Currently amended) The purified nucleic acid of claim 14, wherein said alpha subunit SCN1A nucleic acid encodes:
  - (a) an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 188 which replaces an aspartic acid residue by a valine residue; or
  - (b) an alpha subunit of SCN1A at least 95% identical to the <u>alpha subunit of SCN1A</u> alpha subunits in (a) as set forth in SEQ ID NO:3 and comprising [[said]]a

mutation <u>corresponding to [[at]]</u> amino acid position 188 <u>which replaces an</u> aspartic acid residue by a valine residue.

26.-29. (Canceled)

- (Currently amended) The purified nucleic acid of claim 14, wherein said SCN1A nucleic
  acid fragment [[in (d)]] comprises a GCATTTGACGATATAnucleotide sequence as set forth in
  SEO ID NO:190 or an ATCATATACTTCCTG nucleotide sequence SEO ID NO:192.
- 31. (Canceled)
- 32. (Previously presented) The purified nucleic acid of claim 14, encoding the alpha subunit of SCN1A set forth in SEQ ID NO:3, wherein aspartic acid residue at position 188 is replaced by a value residue.
- 33. (Previously presented) The purified nucleic acid of claim 14, encoding the alpha subunit of SCN1A set forth in SEQ ID NO:3, wherein glutamic acid residue at position 1238 is replaced by an aspartic acid residue.
- 34. (Previously presented) The purified nucleic acid of claim 14, encoding the alpha subunit of SCN1A set forth in SEQ ID NO:3, wherein serine residue at position 1773 is replaced by a tyrosine residue.
- 35.-38. (Canceled)
- (Currently amended) A vector comprising the-sequences any one of the nucleic acids of claim 25.
- (Previously presented) An isolated cell harboring the vector of claim 39.
- 41. (New) A vector comprising any one of the nucleic acids of claim 30.

- 42. (New) An isolated cell harboring the vector of claim 41.
- 43. (New) A vector comprising the nucleic acid of claim 32.
- 44. (New) An isolated cell harboring the vector of claim 43.
- 45. (New) A vector comprising the nucleic acid of claim 33.
- 46. (New) An isolated cell harboring the vector of claim 45.
- 47. (New) A vector comprising the nucleic acid of claim 34.
- 48. (New) An isolated cell harboring the vector of claim 47.
- 49. (New) A purified human alpha subunit of an SCN1A sodium channel nucleic acid comprising a nucleic acid sequence selected from the group consisting of:
  - a nucleic acid sequence encoding an alpha subunit of SCN1A selected from the group consisting of:
    - an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 188 which replaces an aspartic acid residue by a valine residue;
    - (ii) an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 1238 which replaces a glutamic acid residue by an aspartic acid residue;
    - (iii) an alpha subunit of SCN1A as set forth in SEQ ID NO:3, comprising a mutation corresponding to amino acid position 1773 which replaces a serine residue residue by a tyrosine residue; and
    - (iv) an alpha subunit of SCN1A being at least 95% identical to the SCN1A alpha subunits in (i)-(iii) and comprising one of the mutations at amino acid position 188, 1238 or 1773;

- (b) an SCN1A nucleic acid fragment selected from the group consisting of:
  - an amplified segment comprising the nucleic acid sequence from nucleotide 739 to 867 of SEQ ID NO:1 having a mutation at nucleotide 828.
  - an amplified segment comprising the nucleic acid sequence from nucleotide 3970 to 4143 of SEQ ID NO:1 having a mutation at position 3978.
  - (vii) an amplified segment comprising the nucleic acid sequence from nucleotide 5521 to 5747 of SEQ ID NO:1 having a mutation at position 5582; and
- (c) a full-length complement of (a) or (b).
- 50. (New) The nucleic acid of claim 49, wherein said nucleic acid sequence is selected from the group consisting of:
  - (viii) an amplified segment consisting of the nucleic acid sequence from nucleotide 739 to 867 of SEQ ID NO:1 having a mutation at nucleotide 828.
  - (ix) an amplified segment consisting of the nucleic acid sequence from nucleotide 3970 to 4143 of SEQ ID NO:1 having a mutation at position 3978;
  - an amplified segment consisting of the nucleic acid sequence from nucleotide 5521 to 5747 of SEQ ID NO:1 having a mutation at position 5582; and
  - (xi) a full-length complement of (viii)-(x).
- 51. (New) A vector comprising any one of the nucleic acids of claim 49.
- 52. (New) An isolated cell harboring the vector of claim 51.
- 53. (New) A purified human alpha subunit of an SCN1A sodium channel nucleic acid comprising a nucleic acid sequence selected from the group consisting of:

- (a) a nucleic acid sequence encoding an alpha subunit of SCN1A selected from the group consisting of:
  - an alpha subunit of SCN1A as set forth in SEQ ID NO:409, comprising a mutation corresponding to amino acid position 188 which replaces an aspartic acid residue by a valine residue;
  - (ii) an alpha subunit of SCN1A as set forth in SEQ ID NO:410, comprising a mutation corresponding to amino acid position 1238 which replaces a glutamic acid residue by an aspartic acid residue;
  - (iii) an alpha subunit of SCN1A as set forth in SEQ ID NO:411, comprising a mutation corresponding to amino acid position 1773 which replaces a serine residue residue by a tyrosine residue; and
  - (iv) an alpha subunit of SCN1A being at least 95% identical to the SCN1A alpha subunits in (ii)-(iii) and comprising one of the mutations at amino acid position 188, 1238 or 1773; and
- (b) a full-length complement of a).

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